SEQUENCE LISTING

<110> Holloway, James L.

<120> A Method and System for Detecting Near Identities in Large DNA Databases

<130> 99-61

<160> 3

<170> FastSEQ for Windows Version 3.0

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<211> 32

<212> DNA

<213> Artificial Sequence

<220>

<223> Illustrative nucleotide sequence.

<400> 1

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32

<210> 2

<211> 33

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      <213> Artificial Sequence
       <220>
       <223> Illustrative nucleotide sequence.
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       <210> 3
       <211> 50
       <212> DNA
       <213> Artificial Sequence
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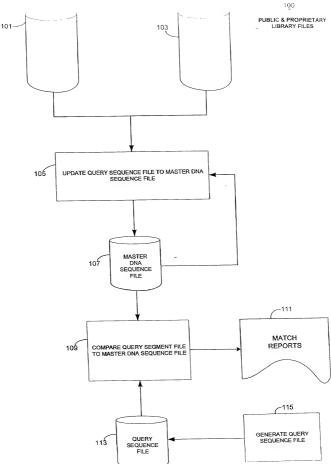
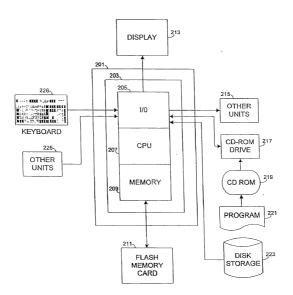
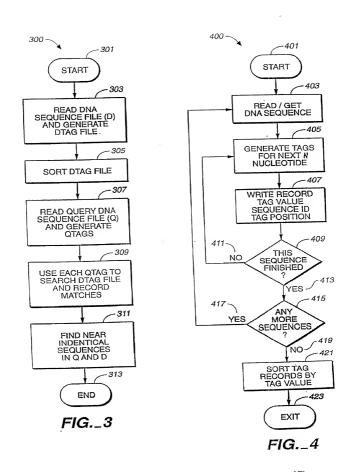


Figure 2
200 Typical General Purpose Computer/





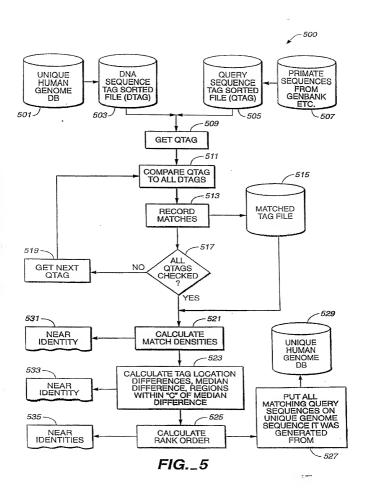


FIGURE 6

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                                               249 1.00
                                     6197
    0.00 3847 933110467
         /vols/pdata/na/genomic/GEN HUMAN.NDB; AQ474914;
635-350 1 289 Z0 753000 557888 558176 9 -1.00 0.00 262-
636-933110467 621
                    627)
         /vols/pdata/na/genomic/GEN HUMAN.NDB; AQ484430;
    346 6 299 ZO 753000 265056 265344 5 -1.00 1.00 262
    933110467
         /vols/pdata/na/genomic/GEN HUMAN.NDB; G34630; 500
            Z0 753000 660416 660832 14 -1.00 0.00 262
    41 457
    933110467
         /vols/pdata/na/genomic/GEN HUMAN.NDB; AQ197006;
     551 41 489 Z0 753000 323936 324384 15 -1.00 0.00 262
     933110467
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                                             14 -1.00 0.00
     2 418
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     6 422
     933110467
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     28 444
     933110467
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     12 429
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     933110467
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FIGURE 6 (continued)

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    390 933110467
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    0.00 390 933110467
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                   15 -1.00 0.00 390 933286163DR
     215264
     /home/disc/holloway/software/genomic_uniq/data/gen_0729
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